

SEQUENCE LISTING

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<110> Allen, Steve
Lee, Jian Ming

<120> Plant Protein Kinases

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<150> 60/092, 438
<151> July 10, 1998

<160> 23

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3/3

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 35 40 45

Ala Thr Gly Lys Glu Phe Ala Cys Lys Ser Ile Leu Lys Xaa Leu Val
 50 55 60

Thr Asp Asp Asp Val Glu Asp Val Arg Arg Glu Ile Gln Ile Met His
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His Leu Ala Gly His Pro Asn Val Ile Ser Ile Arg Gly Ala Tyr Glu
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Asp Ala Val Ala Val
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<213> Oryza sativa

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 35 40 45

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Gly His Gln Thr Pro Gly Val Ala Trp Pro Ser Pro Tyr Pro Ser Gly
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Gly Ala Ser Pro Leu Pro Ala Gly Val Ser Pro Ser Pro Ala Arg Ser
 85 90 95

Thr Pro Arg Arg Phe Phe Lys Arg Pro Phe Pro Pro Pro Ser Pro Ala
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Lys His Ile Lys Ala Thr Leu Ala Lys Arg Leu Gly Gly Lys Pro
 115 120 125

Lys Glu Gly Thr Ile Pro Glu Glu Gly Val Gly Ala Gly Gly Gly
 130 135 140

Gly Gly Gly Ala Ala Asp Gly Ala Glu Thr Glu Arg Pro Leu Asp Lys
 145 150 155 160

Thr Phe Gly Phe Ser Lys Asn Phe Gly Ala Lys Tyr Glu Leu Gly Lys
 165 170 175

Glu Val Gly Arg Gly His Phe Gly His Thr Cys Ser Ala Val Val Lys
180 185 190

Lys Gly Glu Tyr Lys Gly Gln Thr Val Ala Val Lys Ile Ile Ala Lys
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Lys Ile Leu Arg Ala Leu Ser Gly His Asn Asn Leu Val Lys Phe Tyr
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Asp Ala Cys Glu Asp Gly Leu Asn Val Tyr Ile Val Met Glu Leu Cys
245 250 255

Glu Gly Gly Glu Leu Leu Asp Arg Ile Leu Ala Arg Gly Gly Arg Tyr
260 265 270

Thr Glu Glu Asp Ala Lys Ala Ile Val Val Gln Ile Leu Ser Val Val
275 280 285

Ala Phe Cys His Leu Gln Gly Val Val His Arg Asp Leu Lys Pro Glu
290 295 300

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305 310 315 320

Asp Phe Gly Leu Ser Asp Phe Ile Arg Pro Asp Glu Arg Leu Asn Asp
325 330 335

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340 345 350

Tyr Ser Met Glu Ala Asp Ile Trp Ser Ile Gly Val Ile Thr Tyr Ile
355 360 365

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370 375 380

Phe Arg Ser Val Leu Arg Ala Asp Pro Asn Phe Asp Asp Ser Pro Trp
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Pro Thr Val Ser Ala Glu Ala Lys Asp Phe Val Lys Arg Phe Leu Asn
405 410 415

Lys Asp Tyr Arg Lys Arg Met Thr Ala Val Gln Ala Leu Thr His Pro
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Arg Leu Ile Lys Gln Tyr Leu Arg Ala Thr Pro Leu Lys Arg Leu Ala
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Lys Leu Gln Phe Lys Leu Leu Glu Pro Arg Asp Gly Phe Val Ser Leu
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Asp Asn Phe Arg Thr Ala Leu Thr Arg Tyr Leu Thr Asp Ala Met Lys
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Glu Ser Arg Val Leu Glu Phe Leu His Ala Leu Glu Pro Leu Ala Tyr
515 520 525

Arg Arg Met Asp Phe Glu Glu Phe Cys Ala Ala Ala Ile Ser Pro Tyr
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Gln Gln Phe Glu Gln Glu Gly Asn Arg Val Ile Ser Val Glu Glu Leu
565 570 575

Ala Gln Glu Leu Asn Leu Ala Pro Thr His Tyr Ser Ile Val Gln Asp
580 585 590

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Pro Pro Ser Pro Ala Lys His Ile Arg Ala Leu Leu Ala Arg Xaa His
 35 40 45

Gly Ser Val Lys Pro Asn Glu Ala Ser Ile Pro Glu Ala Ser Xaa Cys
 50 55 60

Glu Leu Gly Leu Asp Lys Ser Phe Gly Phe Ala Lys Gln Phe Ser Ala
 65 70 75 80

His Tyr Glu Leu Ser Asp Glu Xaa Gly Arg Gly His Phe Gly Tyr Thr
 85 90 95

Cys Ser Ala Lys Gly Lys Gly Ala Phe Lys Gly Leu Asn Val Ala
 100 105 110

Val Lys Val Ile Pro Lys Ala Lys Met Thr Thr Ala Ile Ala Ile Glu
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Asp Val Arg Arg Glu Val Lys Ile Leu Arg Ala Leu Thr Gly His Lys
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 35 40 45
 Phe Ser Ser Lys Glu Glu Asn Ser Pro Leu Lys Val Ile Asp Phe Gly
 50 55 60
 Leu Ser Asp Phe Val Lys Pro Asp Glu Arg Leu Asn Asp Ile Val Gly
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 <213> Zea mays

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<212> PRT
<213> Zea mays

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35 40 45

Asn Gly Ala Glu Pro Gly His Ile Ile Val Thr Ser Ile Asp Gly Arg
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Asn Gly Gln Ala Lys Gln Thr Ile Ser Tyr Met Ala Glu Arg Val Val
65 70 75 80

Gly His Gly Ser Phe Gly Thr Val Phe Gln Ala Lys Cys Leu Glu Thr
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Gly Glu Thr Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Tyr Lys
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Asn Arg Glu Leu Gln Thr Met Arg Val Leu Asp His Pro Asn Val Val
115 120 125

Ala Leu Lys His Cys Phe Phe Ser Lys Thr Glu Lys Glu Glu Leu Tyr
130 135 140

Leu Asn Leu Val Leu Glu Tyr Val Pro Glu Thr Ala His Arg Val Ile
145 150 155 160

Lys His Tyr Asn Lys Met Asn Gln Arg Met Pro Leu Ile Tyr Ala Lys
165 170 175

Leu Tyr Met Tyr Gln Ile Cys Arg Ala Leu Ala Tyr Ile His Asn Ser
180 185 190

Ile Gly Val Cys His Arg Asp Ile Lys Pro Gln Asn Leu Leu Val Asn
195 200 205

Pro His Thr His Gln Leu Lys Leu Cys Asp Phe Gly Ser Ala Lys Val
210 215 220

Leu Val Lys Gly Glu Pro Asn Ile Ser Tyr Ile Cys Ser Arg Tyr Tyr
225 230 235 240

Arg Ala Pro Glu Leu Ile Phe Gly Ala Thr Glu Tyr Thr Ala Ile
245 250 255

Asp Val Gly Ser Ala Gly Cys Val Leu Ala Glu Leu Leu Gly Gln
260 265 270

Pro Leu Phe Pro Gly Glu Ser Gly Val Asp Gln Leu Val Glu Ile Ile
275 280 285

Lys Val Leu Gly Thr Pro Thr Arg Glu Glu Ile Lys Cys Met Asn Pro
290 295 300

Asn Tyr Thr Glu Phe Lys Phe Pro Gln Ile Lys Ala His Pro Trp His
305 310 315 320

Lys Ile Phe His Lys Arg Met Pro Ala Glu Ala Val Asp Leu Val Ser
325 330 335

Arg Leu Leu Gln Tyr Ser Pro Lys Leu Arg Ser Thr Ala Leu Glu Ala
340 345 350

Leu Val His Pro Phe Phe Asp Glu Leu Arg Asp Pro Asn Thr Arg Leu
355 360 365

Pro Asn Gly Arg Phe Leu Pro Pro Leu Phe Asn Phe Lys Pro His Glu
370 375 380

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<213> Oryza sativa

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35 40 45

Gly Thr Glu Xaa Gly His Ile Ile Val Thr Thr Gly Gly Xaa Asn
50 55 60

Gly Xaa Pro Lys Xaa Thr Val Ser Tyr Met Ala Xaa Arg Ile Val Gly
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Gln Gly Ser Phe Gly Ile Val Phe Gln Ala Lys Phe Trp Arg Gln Gly
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Glycine max

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| aatgaatgg | ttttgagcaa | cagcagttaa | aagagaaáag | ggattcagcg | aagatgacat | 120 |
| cgggttgtgt | ggcaccaact | tcgggttga | gagaaggccag | tggcatgga | gcagcaggtg | 180 |
| ttgatagatt | gccagaggag | atgaacgata | tgaaaattag | ggtatgataga | gaaatggaag | 240 |
| ccacagttgt | tgatggcaac | ggaacggaga | caggacatat | cattgtgact | accattgggg | 300 |
| gtagaaatgg | tcagcccaag | cagactataa | gctacatggc | agagcgtgtt | gtagggcatg | 360 |
| gatcatttgg | agttgtctc | caggctaagt | gcttggaaac | cgtgaaact | gtggctatca | 420 |
| aaaaggttct | tcaagacaag | aggtacaaga | accgggagct | gcaaacaatg | cgccttcttg | 480 |
| accacccaaa | tgtcggtgct | ttgaagcact | gtttctttc | aaccactgaa | aaggatgaac | 540 |
| tataccttaa | tttggttctc | gaatatgttc | ctgaaacagt | taatcgggtt | ataaaacatt | 600 |
| acaacaagtt | taaccaaagg | atgccaetga | tatatgtgaa | actctataca | taccagatct | 660 |
| tttaggcgtt | atcttatatt | catcggtgt | ttggagcttg | ccatcgggat | atcaagcctc | 720 |
| aaaatctatt | ggtcaatcca | cacactcacc | aggttaatt | atgtgacttt | ggaagtgcaa | 780 |
| aggttttgtt | aaaaggcga | ccaaatatat | catacatatg | ttctagatac | tatagagcac | 840 |
| ctgagctcat | atttggcgc | actgaatata | ctacagccat | tgacgtctgg | tctgttggat | 900 |
| gtgttttagc | ttagctgct | tttggacagc | ctctgttccc | tggtagat | ggagttgatc | 960 |
| aacttgttga | gatcatcaag | gttctggca | ctccaacaag | ggaagagatt | aagtgcatt | 1020 |
| accctaatta | tacagaattt | aaatcccac | agattaaagc | acatccatgg | cacaagatct | 1080 |
| tccataagcg | catgcctca | gaggctgtt | atttggtatac | aagactacta | caataactccc | 1140 |
| ctaaacttgc | gtgcacagtt | ttagatgcct | tggacgcacc | cttcctttg | gacgaattcc | 1200 |
| gngatccaaa | tcctcgct | ccaaatggc | cgatccntcc | aacaactatt | aattcaaacc | 1260 |
| catgaactga | aagtgtccaa | ctgagattt | ggaaantgg | tcaaagcatg | caaggaacaa | 1320 |
| tgccgtttct | gcttgtaan | tgtacaaaac | tgaagtgtt | ttcatataga | atgcngctt | 1380 |
| cctcattaaa | gaaattgtgg | accttatgan | tcgttnccgt | aacagttag | | 1429 |

<210> 14
<211> 399
<212> PRT
<213> Glycine max

Det 313
<220>
<221> UNSURE
<222> (391)

<400> 14
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Met Thr Ser Val Gly Val Ala Pro Thr Ser Gly Leu Arg Glu Ala Ser
20 25 30
Gly His Gly Ala Ala Gly Val Asp Arg Leu Pro Glu Glu Met Asn Asp
35 40 45
Met Lys Ile Arg Asp Asp Arg Glu Met Glu Ala Thr Val Val Asp Gly
50 55 60
Asn Gly Thr Glu Thr Gly His Ile Ile Val Thr Thr Ile Gly Gly Arg
65 70 75 80
Asn Gly Gln Pro Lys Gln Thr Ile Ser Tyr Met Ala Glu Arg Val Val
85 90 95
Gly His Gly Ser Phe Gly Val Val Phe Gln Ala Lys Cys Leu Glu Thr
100 105 110
Gly Glu Thr Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Tyr Lys
115 120 125
Asn Arg Glu Leu Gln Thr Met Arg Leu Leu Asp His Pro Asn Val Val
130 135 140
Ala Leu Lys His Cys Phe Phe Ser Thr Thr Glu Lys Asp Glu Leu Tyr
145 150 155 160
Leu Asn Leu Val Leu Glu Tyr Val Pro Glu Thr Val Asn Arg Val Ile
165 170 175
Lys His Tyr Asn Lys Phe Asn Gln Arg Met Pro Leu Ile Tyr Val Lys
180 185 190
Leu Tyr Thr Tyr Gln Ile Phe Arg Ala Leu Ser Tyr Ile His Arg Cys
195 200 205
Ile Gly Val Cys His Arg Asp Ile Lys Pro Gln Asn Leu Leu Val Asn
210 215 220
Pro His Thr His Gln Val Lys Leu Cys Asp Phe Gly Ser Ala Lys Val
225 230 235 240
Leu Val Lys Gly Glu Pro Asn Ile Ser Tyr Ile Cys Ser Arg Tyr Tyr
245 250 255
Arg Ala Pro Glu Leu Ile Phe Gly Ala Thr Glu Tyr Thr Thr Ala Ile
260 265 270
Asp Val Trp Ser Val Gly Cys Val Leu Ala Glu Leu Leu Gly Gln
275 280 285

Pro Leu Phe Pro Gly Glu Ser Gly Val Asp Gln Leu Val Glu Ile Ile
280 295 300

Lys Val Leu Gly Thr Pro Thr Arg Glu Glu Ile Lys Cys Met Asn Pro
 305 310 315 320

Asn Tyr Thr Glu Phe Lys Phe Pro Gln Ile Lys Ala His Pro Trp His
325 330 335

Lys Ile Phe His Lys Arg Met Pro Pro Glu Ala Val Asp Leu Val Ser
340 345 350

Arg Leu Leu Gln Tyr Ser Pro Asn Leu Arg Cys Thr Val Leu Asp Ala
355 360 365

Leu Asp Ala Pro Phe Pro Leu Asp Glu Phe Arg Asp Pro Asn Pro Arg
370 375 380

Leu Pro Asn Gly Pro Ile Xaa Pro Thr Thr Ile Asn Ser Asn Pro
385 390 395

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<210> 15
<211> 1673
<212> DNA
<213> Triticum aestivum
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<220>
<221> unsure
<222> (1349)

<400> 15
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ggcgcggag cccatgctgc tcgacgagca gccccccacc gcagtcgcct gcgagaagaa 120
gcagcaggat ggcgaggcgc cgtatgcgga ggggaacgac gccatgaccg gtcacatcat 180
ctccaccacc atcggcggca agaacggca gcccaagcag acgattagct acatggcgga 240
gcgcgttggc ggcactgggt cgtttggcat cgtcttcag gctaaatgcc tggaaaccgg 300
ggagatgggt ggcattaaga aggtactgca ggacagacgg tacaagaacc gtgagtgca 360
gcttatgcgt tcgatgatcc attccaatgt tgtctccctc aagcactgct tcttctcaac 420
cacaagtaga gatgagotgt tcctgaacct tgcgtatggag tatgtcccg agacgtata 480
ccgcgtgctt aagcaetaca gtaatgccaa ccaggggatg cgccttatct atgtcaagct 540
ttacatgtat cagctttta gagggcttag cttatgttcat actgttccag gagttgcca 600
cagggatgtg aaacccacaaa atgttttgt tgatcctcta acccatcaag tcaagatctg 660
tgactttgga agtgcaaaaag ttctggtacc tggtgaaaccc aacatagcat acatatgctc 720
tcgctactat cgtgctcctg agctcatatt tggtgcaact gaatatacaa cttcaataga 780
catatggtca gctggatgtg ttcttgcaga gctacttctt ggtcagcctc tggttccagg 840
agagactgca gttgatcagc tagtggagat tatcaaggtt cttggtaactc caaccgtga 900
ggaaattcgg tgcataacc ccaactatac cgagttcaagg tttcctcaga ttaaggctca 960
tccttggcac aagattttcc acaagagaat gcccgtgaa gctatagatc ttgcctcccg 1020
ccttctccag tattcaccaa atctacgtt cactgcttt gatgcatgtg cacattcatt 1080
ctttgtatgag ctacgtgagc cgaatgcacg cttggccaaat ggccgcccatt tccctccct 1140
gttcaacttc aaacctgaac tagcgaacgc ctctccagag ctcataaca ggttggttcc 1200
ggaacatgtt cgacggcaaa atggcccaa cttcgcccat gctggagct aaacggggcg 1260
cgccccgcattt gccatattt ttgtttgtcc gccatcatcg aagaatcaat ctctccctta 1320
aatcctgagg agagaccgat caagtgcant gccagtgcacca gtgaaaagaag tacaactatg 1380
taaattacct gaccttggaa gaatcgttgt tgggttgc ggtgcccggcc atgtttaagt 1440
acatggcggc acatgttggt tgagttgtt cttattattt agtaggttaag agcaatgtatg 1500
taggagggtgg agacatatgt taatgttagg tctgtgaccc ttttaagta cattttgtt 1560
atgcttggta gtggtaactgt aatgcggcaat tagctgcctc atgtttgtc cttgtccct 1620
qatqtaaatq tcgtcgccct gcagaaaaaa aaaaaaaaaaaa aaa 1673

Graph
B3

<210> 16
<211> 402
<212> PRT
<213> Triticum aestivum

<400> 16
Met Glu His Pro Ala Pro Ala Pro Glu Pro Met Leu Leu Asp Glu Gln
1 5 10 15
Pro Pro Thr Ala Val Ala Cys Glu Lys Lys Gln Gln Asp Gly Glu Ala
20 25 30
Pro Tyr Ala Glu Gly Asn Asp Ala Met Thr Gly His Ile Ile Ser Thr
35 40 45
Thr Ile Gly Gly Lys Asn Gly Glu Pro Lys Gln Thr Ile Ser Tyr Met
50 55 60
Ala Glu Arg Val Val Gly Thr Gly Ser Phe Gly Ile Val Phe Gln Ala
65 70 75 80
Lys Cys Leu Glu Thr Gly Glu Met Val Gly Ile Lys Lys Val Leu Gln
85 90 95
Asp Arg Arg Tyr Lys Asn Arg Glu Leu Gln Leu Met Arg Ser Met Ile
100 105 110
His Ser Asn Val Val Ser Leu Lys His Cys Phe Phe Ser Thr Thr Ser
115 120 125
Arg Asp Glu Leu Phe Leu Asn Leu Val Met Glu Tyr Val Pro Glu Thr
130 135 140
Leu Tyr Arg Val Leu Lys His Tyr Ser Asn Ala Asn Gln Gly Met Pro
145 150 155 160
Leu Ile Tyr Val Lys Leu Tyr Met Tyr Gln Leu Phe Arg Gly Leu Ala
165 170 175
Tyr Val His Thr Val Pro Gly Val Cys His Arg Asp Val Lys Pro Gln
180 185 190
Asn Val Leu Val Asp Pro Leu Thr His Gln Val Lys Ile Cys Asp Phe
195 200 205
Gly Ser Ala Lys Val Leu Val Pro Gly Glu Pro Asn Ile Ala Tyr Ile
210 215 220
Cys Ser Arg Tyr Tyr Arg Ala Pro Glu Leu Ile Phe Gly Ala Thr Glu
225 230 235 240
Tyr Thr Thr Ser Ile Asp Ile Trp Ser Ala Gly Cys Val Leu Ala Glu
245 250 255
Leu Leu Leu Gly Gln Pro Leu Phe Pro Gly Glu Thr Ala Val Asp Gln
260 265 270
Leu Val Glu Ile Ile Lys Val Leu Gly Thr Pro Thr Arg Glu Glu Ile
275 280 285

Arg Cys Met Asn Pro Asn Tyr Thr Glu Phe Arg Phe Pro Gln Ile Lys
290 295 300

Ala His Pro Trp His Lys Ile Phe His Lys Arg Met Pro Ala Glu Ala
305 310 315 320

Ile Asp Leu Ala Ser Arg Leu Leu Gln Tyr Ser Pro Asn Leu Arg Cys
325 330 335

Thr Ala Leu Asp Ala Cys Ala His Ser Phe Phe Asp Glu Leu Arg Glu
340 345 350

Pro Asn Ala Arg Leu Pro Asn Gly Arg Pro Phe Pro Pro Leu Phe Asn
355 360 365

Phe Lys Pro Glu Leu Ala Asn Ala Ser Pro Glu Leu Ile Asn Arg Leu
370 375 380

Val Pro Glu His Val Arg Arg Gln Asn Gly Pro Asn Phe Ala His Ala
385 390 395 400

Gly Ser

<210> 17

<211> 639

<212> PRT

<213> Zea mays

<400> 17

Met Gly Asn Thr Cys Val Gly Pro Ser Ile Thr Met Asn Gly Phe Phe
1 5 10 15

Gln Ser Val Ser Thr Ala Leu Trp Lys Thr Pro Gln Glu Gly Asp Ala
20 25 30

Leu Pro Ala Ala Ala Asn Gly Pro Gly Gly Pro Ala Gly Ala Gly Ser
35 40 45

Gln Ser Ala Leu Pro Lys Pro Ala Ser Asp Val His His Val Ala Val
50 55 60

Gln Ser Glu Ala Pro Glu Pro Val Lys Ile Ala Ala Tyr His Ser Glu
65 70 75 80

Pro Ala Pro Ala Val Arg Ser Glu Ala Pro Glu Pro Val Lys Ile Ala
85 90 95

Ala Ser His Ser Glu Pro Ala Pro Met Ala Ala Lys Pro Gly Gly Ala
100 105 110

Ala Ala Asn Ala Ser Pro Ser Pro Ser Pro Arg Pro Arg Pro Gln Val
115 120 125

Lys Arg Val Ser Ser Ala Gly Leu Leu Leu Gly Ser Val Leu Arg Arg
130 135 140

Lys Thr Glu Asn Leu Lys Asp Lys Tyr Ser Leu Gly Arg Arg Leu Gly
145 150 155 160

Gln Gly Gln Phe Gly Thr Thr His Leu Cys Val Glu Arg Ala Thr Gly
165 170 175

Lys Glu Leu Ala Cys Lys Ser Ile Leu Lys Arg Lys Leu Gly Ser Asp
180 185 190

Asp Asp Val Glu Asp Val Arg Arg Glu Ile Gln Ile Met His His Leu
195 200 205

Ala Gly His Pro Ser Val Val Gly Ile Arg Gly Ala Tyr Glu Asp Ala
210 215 220

Val Ala Val His Leu Val Met Glu Leu Cys Gly Gly Gly Glu Leu Phe
225 230 235 240

Asp Arg Ile Val Arg Arg Gly His Tyr Thr Glu Arg Lys Ala Ala Glu
245 250 255

Leu Ala Arg Val Ile Val Gly Val Val Glu Ala Cys His Ser Met Gly
260 265 270

Val Met His Arg Asp Leu Lys Pro Glu Asn Phe Leu Phe Ala Asp His
275 280 285

Ser Glu Glu Ala Ala Leu Lys Thr Ile Asp Phe Gly Leu Ser Ile Phe
290 295 300

Phe Arg Pro Gly Gln Ile Phe Thr Asp Val Val Gly Ser Pro Tyr Tyr
305 310 315 320

Val Ala Pro Glu Val Leu Lys Lys Arg Tyr Gly Pro Glu Ala Asp Val
325 330 335

Trp Ser Ala Gly Val Ile Ile Tyr Ile Leu Leu Cys Gly Val Pro Pro
340 345 350

Phe Trp Ala Glu Asn Glu Gln Gly Ile Phe Glu Glu Val Leu His Gly
355 360 365

Arg Leu Asp Phe Glu Ser Glu Pro Trp Pro Ser Ile Ser Asp Gly Ala
370 375 380

Lys Asp Leu Val Arg Arg Met Leu Val Arg Asp Pro Arg Lys Arg Leu
385 390 395 400

Thr Ala His Glu Val Leu Arg His Pro Trp Val Gln Val Gly Gly Val
405 410 415

Ala Pro Asp Arg Pro Leu Asp Ser Ala Val Leu Ser Arg Met Lys Gln
420 425 430

Phe Ser Ala Met Asn Lys Leu Lys Lys Met Ala Leu Arg Val Ile Ala
435 440 445

Glu Asn Leu Ser Glu Asp Glu Ile Ala Gly Leu Arg Glu Met Phe Lys
450 455 460

Met Ile Asp Ala Asp Asn Ser Gly Gln Ile Thr Phe Glu Glu Leu Lys
465 470 475 480

Val Gly Leu Glu Lys Val Gly Ala Asn Leu Gln Glu Ser Glu Ile Tyr
485 490 495

Ala Leu Met Gln Ala Ala Asp Val Asp Asn Asn Gly Thr Ile Asp Tyr
500 505 510

Gly Glu Phe Ile Ala Ala Thr Leu His Leu Asn Lys Val Glu Arg Glu
515 520 525

Asp His Leu Phe Ala Ala Phe Gln Tyr Phe Asp Lys Asp Gly Ser Gly
530 535 540

Tyr Ile Thr Ala Asp Glu Leu Gln Val Ala Cys Glu Glu Phe Gly Leu
545 550 555 560

Gly Asp Val Gln Leu Glu Asp Leu Ile Gly Glu Val Asp Gln Asp Asn
565 570 575

Asp Gly Arg Ile Asp Tyr Asn Glu Phe Val Ala Met Met Gln Lys Pro
580 585 590

Thr Val Gly Gly Ser Arg Arg Arg Pro Ile Cys Arg Thr Ala Ser Ala
595 600 605

Ser Gly Ser Ala Ser Gly Ser Gly Arg Arg Ser Gly Trp Pro Arg Pro
610 615 620

Leu Cys Leu Trp Leu Pro Cys Cys Leu Arg Val Gly Val Asp Asp
625 630 635

<210> 18
<211> 625
<212> PRT
<213> Zea mays

<400> 18
Met Gly Gln Cys Tyr Gly Lys Ala Arg Gly Ala Ser Ser Arg Ala Asp
1 5 10 15

His Asp Ala Asp Pro Ser Gly Ala Gly Ser Val Ala Pro Pro Ser Pro
20 25 30

Leu Pro Ala Asp Gly Ala Pro Leu Pro Ala Thr Pro Arg Arg His Lys
35 40 45

Ser Gly Ser Thr Thr Pro Val His His His Gln Ala Ala Thr Pro Gly
50 55 60

Ala Ala Ala Trp Pro Ser Pro Tyr Pro Ala Gly Gly Ala Ser Pro Leu
65 70 75 80

Pro Ala Gly Val Ser Pro Ser Pro Ala Arg Ser Thr Pro Arg Arg Phe
85 90 95

Phe Lys Arg Pro Phe Pro Pro Ser Pro Ala Lys His Ile Lys Ala
100 105 110

Thr Leu Ala Lys Arg Leu Gly Gly Lys Pro Lys Glu Gly Thr Ile
115 120 125

Pro Glu Glu Gly Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala
130 135 140

Gly Ala Ala Val Gly Ala Ala Asp Ser Ala Glu Ala Asp Arg Pro Leu
145 150 155 160

Asp Lys Thr Phe Gly Phe Ala Lys Asn Phe Gly Ala Lys Tyr Asp Leu
165 170 175

Cat
7/13

Gly Lys Glu Val Gly Arg Gly His Phe Gly His Thr Cys Ser Ala Val
180 185 190

Val Lys Lys Gly Glu His Lys Gly His Thr Val Ala Val Lys Ile Ile
195 200 205

Ser Lys Ala Lys Met Thr Thr Ala Ile Ser Ile Glu Asp Val Arg Arg
210 215 220

Glu Val Lys Ile Leu Lys Ala Leu Ser Gly His Asp Asp Leu Val Arg
225 230 235 240

Phe Tyr Asp Ala Cys Glu Asp Ala Leu Asn Val Tyr Ile Val Met Glu
245 250 255

Leu Cys Glu Gly Glu Leu Leu Asp Arg Ile Leu Ala Arg Gly Gly
260 265 270

Arg Tyr Thr Glu Glu Asp Ala Lys Ala Ile Ile Val Gln Ile Leu Ser
275 280 285

Val Val Ala Phe Cys His Leu Gln Gly Val Val His Arg Asp Leu Lys
290 295 300

Pro Glu Asn Phe Leu Phe Thr Thr Arg Asp Glu Ser Ala Pro Met Lys
305 310 315 320

Leu Ile Asp Phe Gly Leu Ser Asp Phe Ile Arg Pro Asp Glu Arg Leu
325 330 335

Asn Asp Ile Val Gly Ser Ala Tyr Tyr Val Ala Pro Glu Val Leu His
340 345 350

Arg Ser Tyr Ser Met Glu Ala Asp Ile Trp Ser Ile Gly Val Ile Thr
355 360 365

Tyr Ile Leu Leu Cys Gly Ser Arg Pro Phe Trp Ala Arg Thr Glu Ser
370 375 380

Gly Ile Phe Arg Ser Val Leu Arg Ala Asp Pro Asn Phe Asp Asp Ser
385 390 395 400

Pro Trp Pro Ser Val Ser Ala Glu Ala Lys Asp Phe Val Lys Arg Phe
405 410 415

Leu Asn Lys Asp Tyr Arg Lys Arg Met Thr Ala Val Gln Ala Leu Thr
420 425 430

His Pro Trp Leu Arg Asp Glu Gln Arg Gln Ile Pro Leu Asp Ile Leu
435 440 445

450 Ile Phe Arg Leu Val Lys Gln Tyr Leu Arg Ala Thr Pro Leu Lys Arg
455 460

465 Leu Ala Leu Lys Ala Leu Ser Lys Ala Leu Ser Glu Asp Glu Leu Leu
470 475 480

485 Tyr Leu Arg Leu Gln Phe Lys Leu Leu Glu Pro Arg Asp Gly Phe Val
490 495

500 Ser Leu Asp Asn Phe Arg Thr Ala Leu Thr Arg Tyr Ser Thr Asp Ala
505 510

515 Met Arg Glu Ser Arg Val Leu Glu Phe Gln His Ala Leu Glu Pro Leu
520 525

530 Ala Tyr Arg Lys Met Asp Phe Glu Glu Phe Cys Ala Ala Ala Ile Ser
535 540

545 Pro Tyr Gln Leu Glu Ala Leu Glu Arg Trp Glu Glu Ile Ala Gly Thr
550 555 560

565 Ala Phe Gln His Phe Glu Gln Glu Gly Asn Arg Val Ile Ser Val Glu
570 575

580 Glu Leu Ala Gln Glu Leu Asn Leu Ala Pro Thr His Tyr Ser Ile Val
585 590

595 Gln Asp Trp Ile Arg Lys Ser Asp Gly Lys Leu Asn Phe Leu Gly Phe
600 605

610 Thr Lys Phe Leu His Gly Val Thr Ile Arg Gly Ser Asn Thr Arg Arg
615 620

625 His

<210> 19
<211> 576
<212> PRT
<213> *Arabidopsis thaliana*

<400> 19
1 Met Gly Ile Cys His Gly Lys Pro Val Glu Gln Gln Ser Lys Ser Leu
5 10 15

20 Pro Val Ser Gly Glu Thr Asn Glu Ala Pro Thr Asn Ser Gln Pro Pro
25 30

35 Ala Lys Ser Ser Gly Phe Pro Phe Tyr Ser Pro Ser Pro Val Pro Ser
40 45

50 Leu Phe Lys Ser Ser Pro Ser Val Ser Ser Ser Val Ser Ser Thr Pro
55 60

65 Leu Arg Ile Phe Lys Arg Pro Phe Pro Pro Ser Pro Ala Lys His
70 75 80

85 Ile Arg Ala Phe Leu Ala Arg Arg Tyr Gly Ser Val Lys Pro Asn Glu
90 95

Val Ser Ile Pro Glu Gly Lys Glu Cys Glu Ile Gly Leu Asp Lys Ser
 100 105 110

Phe Gly Phe Ser Lys Gln Phe Ala Ser His Tyr Glu Ile Asp Gly Glu
 115 120 125

Val Gly Arg Gly His Phe Gly Tyr Thr Cys Ser Ala Lys Gly Lys Lys
 130 135 140

Gly Ser Leu Lys Gly Gln Glu Val Ala Val Lys Val Ile Pro Lys Ser
 145 150 155 160

Lys Met Thr Thr Ala Ile Ala Ile Glu Asp Val Ser Arg Glu Val Lys
 165 170 175

Met Leu Arg Ala Leu Thr Gly His Lys Asn Leu Val Gln Phe Tyr Asp
 180 185 190

Ala Phe Glu Asp Asp Glu Asn Val Tyr Ile Val Met Glu Leu Cys Lys
 195 200 205

Gly Gly Glu Leu Leu Asp Lys Ile Leu Gln Arg Gly Gly Lys Tyr Ser
 210 215 220

Glu Asp Asp Ala Lys Lys Val Met Val Gln Ile Leu Ser Val Val Ala
 225 230 235 240

Tyr Cys His Leu Gln Gly Val Val His Arg Asp Leu Lys Pro Glu Asn
 245 250 255

Phe Leu Phe Ser Thr Lys Asp Glu Thr Ser Pro Leu Lys Ala Ile Asp
 260 265 270

Phe Gly Leu Ser Asp Tyr Val Lys Pro Asp Glu Arg Leu Asn Asp Ile
 275 280 285

Val Gly Ser Ala Tyr Tyr Val Ala Pro Glu Val Leu His Arg Thr Tyr
 290 295 300

Gly Thr Glu Ala Asp Met Trp Ser Ile Gly Val Ile Ala Tyr Ile Leu
 305 310 315 320

Leu Cys Gly Ser Arg Pro Phe Trp Ala Arg Thr Glu Ser Gly Ile Phe
 325 330 335

Arg Ala Val Leu Lys Ala Glu Pro Asn Phe Glu Glu Ala Pro Trp Pro
 340 345 350

Ser Leu Ser Pro Glu Ala Val Asp Phe Val Lys Arg Leu Leu Asn Lys
 355 360 365

Asp Tyr Arg Lys Arg Leu Thr Ala Ala Gln Ala Leu Cys His Pro Trp
 370 375 380

Leu Val Gly Ser His Glu Leu Lys Ile Pro Ser Asp Met Ile Ile Tyr
 385 390 395 400

Lys Leu Val Lys Val Tyr Ile Met Ser Thr Ser Leu Arg Lys Ser Ala
 405 410 415

Leu Ala Ala Leu Ala Lys Thr Leu Thr Val Pro Gln Leu Ala Tyr Leu
 420 425 430

Arg Glu Gln Phe Thr Leu Leu Gly Pro Ser Lys Asn Gly Tyr Ile Ser
 435 440 445

Met Gln Asn Tyr Lys Thr Ala Ile Leu Lys Ser Ser Thr Asp Ala Met
 450 455 460

Lys Asp Ser Arg Val Phe Asp Phe Val His Met Ile Ser Cys Leu Gln
 465 470 475 480

Tyr Lys Lys Leu Asp Phe Glu Glu Phe Cys Ala Ser Ala Leu Ser Val
 485 490 495

Tyr Gln Leu Glu Ala Met Glu Thr Trp Glu Gln His Ala Arg Arg Ala
 500 505 510

Tyr Glu Leu Phe Glu Lys Asp Gly Asn Arg Pro Ile Met Ile Glu Glu
 515 520 525

Leu Ala Ser Glu Leu Gly Leu Gly Pro Ser Val Pro Val His Val Val
 530 535 540

Leu Gln Asp Trp Ile Arg His Ser Asp Gly Lys Leu Ser Phe Leu Gly
 545 550 555 560

Phe Val Arg Leu Leu His Gly Val Ser Ser Arg Thr Leu Gln Lys Ala
 565 570 575

<210> 20
 <211> 405
 <212> PRT
 <213> *Arabidopsis thaliana*

<400> 20
 Met Ala Ser Val Gly Ile Ala Pro Asn Pro Gly Ala Arg Asp Ser Thr
 1 5 10 15

Gly Val Asp Lys Leu Pro Glu Glu Met Asn Asp Met Lys Ile Arg Asp
 20 25 30

Asp Lys Glu Met Glu Ala Thr Val Val Asp Gly Asn Gly Thr Glu Thr
 35 40 45

Gly His Ile Ile Val Thr Thr Ile Gly Gly Arg Asn Gly Gln Pro Lys
 50 55 60

Gln Thr Ile Ser Tyr Met Ala Glu Arg Val Val Gly His Gly Ser Phe
 65 70 75 80

Gly Val Val Phe Gln Ala Lys Cys Leu Glu Thr Gly Glu Thr Val Ala
 85 90 95

Ile Lys Lys Val Leu Gln Asp Arg Arg Tyr Lys Asn Arg Glu Leu Gln
 100 105 110

Thr Met Arg Leu Leu Asp His Pro Asn Val Val Ser Leu Lys His Cys
 115 120 125

Phe Phe Ser Thr Thr Glu Lys Asp Glu Leu Tyr Leu Asn Leu Val Leu
 130 135 140

Glu Tyr Val Pro Glu Thr Val His Arg Val Ile Lys His Tyr Asn Lys
 145 150 155 160

Cat
B13

Leu Asn Gln Arg Met Pro Leu Ile Tyr Val Lys Leu Tyr Thr Tyr Gln
 165 170 175

Ile Phe Arg Ala Leu Ser Tyr Ile His Arg Cys Ile Gly Val Cys His
 180 185 190

Arg Asp Ile Lys Pro Gln Asn Leu Leu Val Asn Pro His Thr His Gln
 195 200 205

Val Lys Leu Cys Asp Phe Gly Ser Ala Lys Val Leu Val Lys Gly Glu
 210 215 220

Pro Asn Ile Ser Tyr Ile Cys Ser Arg Tyr Tyr Arg Ala Pro Glu Leu
 225 230 235 240

Ile Phe Gly Ala Thr Glu Tyr Thr Thr Ala Ile Asp Val Trp Ser Ala
 245 250 255

Gly Cys Val Leu Ala Glu Leu Leu Gly Gln Pro Leu Phe Pro Gly
 260 265 270

Glu Ser Gly Val Asp Gln Leu Val His Ile Ile Lys Val Leu Gly Thr
 275 280 285

Pro Thr Arg Glu Glu Ile Lys Cys Met Asn Pro Asn Tyr Thr Glu Phe
 290 295 300

Lys Phe Pro Gln Ile Lys Ala His Pro Trp His Lys Ile Phe His Lys
 305 310 315 320

Arg Met Pro Pro Glu Ala Val Asp Leu Val Ser Arg Leu Leu Gln Tyr
 325 330 335

Ser Pro Asn Leu Arg Ser Ala Ala Leu Asp Thr Leu Val His Pro Phe
 340 345 350

Phe Asp Glu Leu Arg Asp Pro Asn Ala Arg Leu Pro Asn Gly Arg Phe
 355 360 365

Leu Pro Pro Ala Phe His Phe Lys Pro His Glu Leu Lys Gly Val Pro
 370 375 380

Leu Glu Met Val Ala Lys Leu Val Pro Glu His Ala Arg Lys Gln Cys
 385 390 395 400

Pro Trp Leu Gly Leu
 405

<210> 21
 <211> 412
 <212> PRT
 <213> *Medicago sativa*

<400> 21

Met Met Ala Ser Gly Gly Val Ala Pro Ala Ser Gly Phe Ile Asp Lys
1 5 10 15

Asn Ala Ser Ser Val Gly Val Glu Lys Leu Pro Glu Glu Met Asn Asp
20 25 30

Met Lys Ile Arg Asp Asp Lys Glu Met Glu Ala Ala Thr Ile Val Asp
35 40 45

Gly Asn Gly Thr Glu Thr Gly His Ile Ile Val Thr Thr Ile Gly Gly
50 55 60

Lys Asn Gly Gln Pro Lys Gln Thr Ile Ser Tyr Met Ala Glu Arg Val
65 70 75 80

Val Gly His Gly Ser Phe Gly Val Val Phe Gln Ala Lys Cys Leu Glu
85 90 95

Thr Gly Glu Thr Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Tyr
100 105 110

Lys Asn Arg Glu Leu Gln Thr Met Arg Leu Leu Asp His Pro Asn Val
115 120 125

Val Ser Leu Lys His Cys Phe Phe Ser Thr Thr Glu Lys Asp Glu Leu
130 135 140

Tyr Leu Asn Leu Val Leu Glu Tyr Val Pro Glu Thr Val Ser Arg Val
145 150 155 160

Ile Arg His Tyr Asn Lys Met Asn Gln Arg Met Pro Met Ile Tyr Val
165 170 175

Lys Leu Tyr Ser Tyr Gln Ile Cys Arg Ala Leu Ala Tyr Ile His Asn
180 185 190

Ser Ile Gly Val Cys His Arg Asp Ile Lys Pro Gln Asn Leu Leu Val
195 200 205

Asn Pro His Thr His Gln Leu Lys Ile Cys Asp Phe Gly Ser Ala Lys
210 215 220

Val Leu Val Lys Gly Glu Pro Asn Ile Ser Tyr Ile Cys Ser Arg Tyr
225 230 235 240

Tyr Arg Ala Pro Glu Leu Ile Phe Gly Ala Thr Glu Tyr Thr Thr Ala
245 250 255

Ile Asp Ile Trp Ser Ala Gly Cys Val Leu Gly Glu Leu Leu Gly
260 265 270

Gln Pro Leu Phe Pro Gly Glu Ser Gly Val Asp Gln Leu Val Glu Ile
275 280 285

Ile Lys Val Leu Gly Thr Pro Thr Arg Glu Glu Ile Lys Cys Met Asn
290 295 300

Pro Asn Tyr Thr Glu Phe Lys Phe Pro Gln Ile Lys Ala His Pro Trp
305 310 315 320

His Lys Ile Phe His Lys Arg Met Pro Pro Glu Ala Val Asp Leu Val
325 330 335

Ser Arg Leu Leu Gln Tyr Ser Pro Asn Leu Arg Ser Thr Ala Leu Glu
340 345 350

Ala Leu Val His Pro Phe Tyr Asp Asp Val Arg Asp Pro Asn Thr Arg
355 360 365

Leu Pro Asn Gly Arg Phe Leu Pro Pro Leu Phe Asn Phe Lys Val Asn
370 375 380

Glu Leu Lys Gly Val Pro Ala Glu Met Leu Val Lys Leu Val Pro Pro
385 390 395 400

His Ala Arg Lys Gln Cys Ala Leu Phe Gly Ser Ser
405 410

<210> 22

<211> 411

<212> PRT

<213> *Medicago sativa*

<400> 22

Met Ala Ser Val Gly Val Ala Pro Thr Ser Gly Phe Arg Glu Val Leu
1 5 10 15

Gly Asp Gly Glu Ile Gly Val Asp Asp Ile Leu Pro Glu Glu Met Ser
20 25 30

Asp Met Lys Ile Arg Asp Asp Arg Glu Met Glu Ala Thr Val Val Asp
35 40 45

Gly Asn Gly Thr Glu Thr Gly His Ile Ile Val Thr Thr Ile Gly Gly
50 55 60

Arg Asn Gly Gln Pro Lys Gln Thr Ile Ser Tyr Met Ala Glu Arg Val
65 70 75 80

Val Gly His Gly Ser Phe Gly Val Val Phe Gln Ala Lys Cys Leu Glu
85 90 95

Thr Gly Glu Thr Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Tyr
100 105 110

Lys Asn Arg Glu Leu Gln Thr Met Arg Leu Leu Asp His Pro Asn Val
115 120 125

Val Ser Leu Lys His Cys Phe Phe Ser Thr Thr Glu Lys Asp Glu Leu
130 135 140

Tyr Leu Asn Leu Val Leu Glu Tyr Val Pro Glu Thr Val His Arg Val
145 150 155 160

Ile Lys His Tyr Ser Lys Leu Asn Gln Arg Met Pro Met Ile Tyr Val
165 170 175

Lys Leu Tyr Thr Tyr Gln Ile Phe Arg Ala Leu Ser Tyr Ile His Arg
180 185 190

Cys Ile Gly Val Cys His Arg Asp Ile Lys Pro Gln Asn Leu Leu Val
195 200 205

Asn Pro His Thr His Gln Val Lys Leu Cys Asp Phe Gly Ser Ala Lys
210 215 220

Val Leu Val Lys Gly Glu Pro Asn Ile Ser Tyr Ile Cys Ser Arg Tyr
225 230 235 240

Tyr Arg Ala Pro Glu Leu Ile Phe Gly Ala Thr Glu Tyr Thr Thr Ala
245 250 255

Ile Asp Val Trp Ser Val Gly Cys Val Leu Ala Glu Leu Leu Gly
260 265 270

Gln Pro Leu Phe Pro Gly Glu Arg Gly Val Asp Gln Leu Val Glu Ile
275 280 285

Ile Lys Val Leu Gly Thr Pro Thr Arg Glu Glu Ile Lys Cys Met Asn
290 295 300

Pro Asn Tyr Thr Glu Phe Lys Phe Pro Gln Ile Lys Ala His Pro Trp
305 310 315 320

His Lys Ile Phe His Lys Arg Met Pro Ala Glu Ala Val Asp Leu Val
325 330 335

Ser Arg Leu Leu Gln Tyr Ser Pro Asn Leu Arg Cys Gln Ala Leu Asp
340 345 350

Cys Leu Thr His Pro Phe Phe Asp Glu Leu Arg Asp Pro Asn Ala Arg
355 360 365

Leu Pro Thr Gly Arg Phe Leu Pro Pro Leu Phe Asn Phe Lys Pro His
370 375 380

Glu Leu Lys Gly Val Pro Val Glu Thr Leu Met Lys Leu Val Pro Glu
385 390 395 400

His Ala Arg Lys Gln Cys Pro Phe Leu Gly Leu
405 410

<210> 23

<211> 407

<212> PRT

<213> *Arabidopsis thaliana*

<400> 23

Met Ala Ser Leu Pro Leu Gly Pro Gln Pro His Ala Leu Ala Pro Pro
1 5 10 15

Leu Gln Leu His Asp Gly Asp Ala Leu Lys Arg Arg Pro Glu Leu Asp
20 25 30

Ser Asp Lys Glu Met Ser Ala Ala Val Ile Glu Gly Asn Asp Ala Val
35 40 45

Thr Gly His Ile Ile Ser Thr Thr Ile Gly Gly Lys Asn Gly Glu Pro
50 55 60

Lys Gln Thr Ile Ser Tyr Met Ala Glu Arg Val Val Gly Thr Gly Ser
 65 70 75 80

Phe Gly Ile Val Phe Gln Ala Lys Cys Leu Glu Thr Gly Glu Ser Val
 85 90 95

Ala Ile Lys Lys Val Leu Gln Asp Arg Arg Tyr Lys Asn Arg Glu Leu
 100 105 110

Gln Leu Met Arg Pro Met Asp His Pro Asn Val Ile Ser Leu Lys His
 115 120 125

Cys Phe Phe Ser Thr Thr Ser Arg Asp Glu Leu Phe Leu Asn Leu Val
 130 135 140

Met Glu Tyr Val Pro Glu Thr Leu Tyr Arg Val Leu Arg His Tyr Thr
 145 150 155 160

Ser Ser Asn Gln Arg Met Pro Ile Phe Tyr Val Lys Leu Tyr Thr Tyr
 165 170 175

Gln Ile Phe Arg Gly Leu Ala Tyr Ile His Thr Val Pro Gly Val Cys
 180 185 190

His Arg Asp Val Lys Pro Gln Asn Leu Leu Val Asp Pro Leu Thr His
 195 200 205

Gln Val Lys Leu Cys Asp Phe Gly Ser Ala Lys Val Leu Val Lys Gly
 210 215 220

Glu Pro Asn Ile Ser Tyr Ile Cys Ser Arg Tyr Tyr Arg Ala Pro Glu
 225 230 235 240

Leu Ile Phe Gly Ala Thr Glu Tyr Thr Ala Ser Ile Asp Ile Trp Ser
 245 250 255

Ala Gly Cys Val Leu Ala Glu Leu Leu Leu Gly Gln Pro Leu Phe Pro
 260 265 270

Gly Glu Asn Ser Val Asp Gln Leu Val Glu Ile Ile Lys Val Leu Gly
 275 280 285

Thr Pro Thr Arg Glu Glu Ile Arg Cys Met Asn Pro Asn Tyr Thr Asp
 290 295 300

Phe Arg Phe Pro Gln Ile Lys Ala His Pro Trp His Lys Val Phe His
 305 310 315 320

Lys Arg Met Pro Pro Glu Ala Ile Asp Leu Ala Ser Arg Leu Leu Gln
 325 330 335

Tyr Ser Pro Ser Leu Arg Cys Thr Ala Leu Glu Ala Cys Ala His Pro
 340 345 350

Phe Phe Asn Glu Leu Arg Glu Pro Asn Ala Arg Leu Pro Asn Gly Arg
 355 360 365

Pro Leu Pro Pro Leu Phe Asn Phe Lys Gln Glu Leu Gly Gly Ala Ser
 370 375 380

Met Glu Leu Ile Asn Arg Leu Ile Pro Glu His Val Arg Arg Gln Met
385 390 395 400

Ser Thr Gly Leu Gln Asn Ser
405

Cost
B13

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